

SEQUENCE LISTING

<110> Newell, Martha K.

<120> METHODS AND PRODUCTS RELATED TO
METABOLIC INTERACTIONS IN DISEASE

<130> V0139/7028/HK

<150> U.S. 60/082,250

<151> 1998-04-17

<150> U.S. 60/094,519

<151> 1998-07-29

<150> U.S. 60/101,580

<151> 1998-09-24

<160> 13

<170> FastSEQ for Windows Version 3.0

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<211> 1491

<212> DNA

<213> Homo Sapiens

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cctaagcatc	tgaagccatg	ggccacacac	ggaggcgagg	aacatcacca	tccaagtgtc	360
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Tyr	Trp	Gln	Lys	Glu	Lys	Lys	Met	Val	Leu	Thr	Met	Met	Ser	Gly	Asp
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Met	Asn	Ile	Trp	Pro	Glu	Tyr	Lys	Asn	Arg	Thr	Ile	Phe	Asp	Ile	Thr
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Asn	Asn	Leu	Ser	Ile	Val	Ile	Leu	Ala	Leu	Arg	Pro	Ser	Asp	Glu	Gly
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Thr	Tyr	Glu	Cys	Val	Val	Leu	Lys	Tyr	Glu	Lys	Asp	Ala	Phe	Lys	Arg
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Glu	His	Leu	Ala	Glu	Val	Thr	Leu	Ser	Val	Lys	Ala	Asp	Phe	Pro	Thr
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Pro	Ser	Ile	Ser	Asp	Phe	Glu	Ile	Pro	Thr	Ser	Asn	Ile	Arg	Arg	Ile
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Ile	Cys	Ser	Thr	Ser	Gly	Gly	Phe	Pro	Glu	Pro	His	Leu	Ser	Trp	Leu
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Thr	Thr	Asn	His	Ser	Phe	Met	Cys	Leu	Ile	Lys	Tyr	Gly	His	Leu	Arg
	210					215						220			
Val	Asn	Gln	Thr	Phe	Asn	Trp	Asn	Thr	Thr	Lys	Gln	Glu	His	Phe	Pro
225					230					235					240
Asp	Asn	Leu	Leu	Pro	Ser	Trp	Ala	Ile	Thr	Leu	Ile	Ser	Val	Asn	Gly
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Ile	Phe	Val	Ile	Cys	Cys	Leu	Thr	Tyr	Cys	Phe	Ala	Pro	Arg	Cys	Arg
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cagtggacag	gcatttgtga	cagcactatg	ggactgagta	acattctctt	tgtgatggcc	180
ttcctgctct	ctggtgctgc	tcctctgaag	attcaagctt	atttcaatga	gactgcagac	240
ctgccatgcc	aatttgcaaa	ctctcaaaac	caaagcctga	gtgagctagt	agtattttgg	300
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 260 265 270
 Ser Tyr Lys Cys Gly Thr Asn Thr Met Glu Arg Glu Glu Ser Glu Gln
 275 280 285
 Thr Lys Lys Arg Glu Lys Ile His Ile Pro Glu Arg Ser Asp Glu Ala
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 Gln Arg Val Phe Lys Ser Ser Lys Thr Ser Ser Cys Asp Lys Ser Asp
 305 310 315 320
 Thr Cys Phe

<210> 5
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 <213> Homo Sapiens

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 35 40 45
 Thr Ser Ser Val Ile Arg Tyr Lys Gly Val Leu Gly Thr Ile Thr Ala
 50 55 60
 Val Val Lys Thr Glu Gly Arg Met Lys Leu Tyr Ser Gly Leu Pro Ala
 65 70 75 80
 Gly Leu Gln Arg Gln Ile Ser Ser Ala Ser Leu Arg Ile Gly Leu Tyr
 85 90 95
 Asp Thr Val Gln Glu Phe Leu Thr Ala Gly Lys Glu Thr Ala Pro Ser

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Leu Gly Ser Lys Ile Leu Ala Gly Leu Thr Thr Gly Gly Val Ala Val					
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Phe Ile Gly Gln Pro Thr Glu Val Val Lys Val Arg Leu Gln Ala Gln					
130		135		140	
Ser His Leu His Gly Ile Lys Pro Arg Tyr Thr Gly Thr Tyr Asn Ala					
145		150		155	160
Tyr Arg Ile Ile Ala Thr Thr Glu Gly Leu Thr Gly Leu Trp Lys Gly					
	165		170		175
Thr Thr Pro Asn Leu Met Arg Ser Val Ile Ile Asn Cys Thr Glu Leu					
	180		185		190
Val Thr Tyr Asp Leu Met Lys Glu Ala Phe Val Lys Asn Asn Ile Leu					
	195		200		205
Ala Asp Asp Val Pro Cys His Leu Val Ser Ala Leu Ile Ala Gly Phe					
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Cys Ala Thr Ala Met Ser Ser Pro Val Asp Val Val Lys Thr Arg Phe					
225		230		235	240
Ile Asn Ser Pro Pro Gly Gln Tyr Lys Ser Val Pro Asn Cys Ala Met					
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Lys Val Phe Thr Asn Glu Gly Pro Thr Ala Phe Phe Lys Gly Leu Val					
	260		265		270
Pro Ser Phe Leu Arg Leu Gly Ser Trp Asn Val Ile Met Phe Val Cys					
	275		280		285
Phe Glu Gln Leu Lys Arg Glu Leu Ser Lys Ser Arg Gln Thr Met Asp					
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Cys Ala Thr					
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<400> 8

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			20					25					30		
Pro	Leu	Asp	Thr	Ala	Lys	Val	Arg	Leu	Gln	Ile	Gln	Gly	Glu	Ser	Gln
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Thr	Ile	Leu	Thr	Met	Val	Arg	Thr	Glu	Gly	Pro	Arg	Ser	Leu	Tyr	Asn
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Gly	Leu	Val	Ala	Gly	Leu	Gln	Arg	Gln	Met	Ser	Phe	Ala	Ser	Val	Arg
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Ile	Gly	Leu	Tyr	Asp	Ser	Val	Lys	Gln	Phe	Tyr	Thr	Lys	Gly	Ser	Glu
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			165						170					175	
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		180						185					190		
Glu	Leu	Val	Thr	Tyr	Asp	Leu	Ile	Lys	Asp	Ala	Leu	Leu	Lys	Ala	Asn
	195					200						205			
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Arg	Tyr	Met	Asn	Ser	Ala	Leu	Gly	Gln	Tyr	Ser	Ser	Ala	Gly	His	Cys
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Ala	Leu	Thr	Met	Leu	Gln	Lys	Glu	Gly	Pro	Arg	Ala	Phe	Tyr	Lys	Gly
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Phe	Met	Pro	Ser	Phe	Leu	Arg	Leu	Gly	Ser	Trp	Asn	Val	Val	Met	Phe
	275					280						285			
Val	Thr	Tyr	Glu	Gln	Leu	Lys	Arg	Ala	Leu	Met	Ala	Ala	Cys	Thr	Ser
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 <213> Homo Sapiens

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			20					25					30		
Pro	Leu	Asp	Thr	Ala	Lys	Val	Arg	Leu	Gln	Ile	Gln	Gly	Glu	Asn	Gln
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Ala	Val	Gln	Thr	Ala	Arg	Leu	Val	Gln	Tyr	Arg	Gly	Val	Leu	Gly	Thr
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Ile	Leu	Thr	Met	Val	Arg	Thr	Glu	Gly	Pro	Cys	Ser	Pro	Tyr	Asn	Gly
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Met	Ala	Val	Thr	Cys	Ala	Gln	Pro	Thr	Asp	Val	Val	Lys	Val	Arg	Phe
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Gln	Ala	Ser	Ile	His	Leu	Gly	Pro	Ser	Arg	Ser	Asp	Arg	Lys	Tyr	Ser
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Gly	Leu	Trp	Lys	Gly	Thr	Leu	Pro	Asn	Ile	Met	Arg	Asn	Ala	Ile	Val
			180					185					190		
Asn	Cys	Ala	Glu	Val	Val	Thr	Tyr	Asp	Ile	Leu	Lys	Glu	Lys	Leu	Leu
			195				200						205		
Asp	Tyr	His	Leu	Leu	Thr	Asp	Asn	Phe	Pro	Cys	His	Phe	Val	Ser	Ala
	210					215						220			
Phe	Gly	Ala	Gly	Phe	Cys	Ala	Thr	Val	Val	Ala	Ser	Pro	Val	Asp	Val
225					230					235					240
Val	Lys	Thr	Arg	Tyr	Met	Asn	Ser	Pro	Pro	Gly	Gln	Tyr	Phe	Ser	Pro
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